



[illegible]

1 RESULTS 2  
 2 DTG 08 5420 4740 4750  
 3 SEPTEMBER 4, APPLICATION US76084,203740  
 4 Patent No. 4,451,753  
 5 GENERAL INFORMATION:  
 6 APPLICANT: Lombard Link, Doyle  
 7 APPLICANT: Stooler, Frank R  
 8 APPLICANT: Schaefer, Gerald J  
 9 APPLICANT: Brown, Stella P  
 10 APPLICANT: Johnson, Lincoln V  
 11 APPLICANT: Redington, Theodore R  
 12 TITLE OR INVENTION: RETINAL PROMINENT EYE ILLIUM RECEIVED, REFLECTED FACTOR  
 13 FILE REFERENCE: 2026 420484  
 14 CURRENT APPLICATOR NUMBER: 045067750, 5741  
 15 CURRENT FILING DATE: 1995 08 29  
 16 PRIOR APPLICATION NUMBER: 084779,719  
 17 PRIOR FILING DATE: 1995 01 25  
 18 PRIOR APPLICATION NUMBER: 084779,979  
 19 PRIOR FILING DATE: 1994 07 25  
 20 PRIOR APPLICATION NUMBER: 072904,215  
 21 PRIOR FILING DATE: 1992 06 04  
 22 PRIOR APPLICATION NUMBER: 072952,796  
 23 PRIOR FILING DATE: 1992 09 24  
 24 NUMBER OF PATENTS: 41  
 25 SOFTWARE: Patented Ver. 2.1  
 26 SEQ ID NO 4  
 27 LENGTH: 14581  
 28 TYPE: DNA  
 29 ORGANISM: HUMAN  
 30 FEATURE:  
 31 OTHER INFORMATION: MENA: 66884; EXON: 66884-700; EXIN 11564 11675;  
 32 OTHER INFORMATION: EXON: 14569, 14581; INTRON: 6701, 11564; INTRON:  
 33 5200 3740 4  
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RESID. 3  
US-08-667-841A-43/c  
Sequence 43, Application US/08667841A  
Patent No. 6319687

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1 GENERAL INFORMATION:
2 APPLICANT: Chadoe, Gerald J.; Kaitanoze,
3 APPLICANT: Taniato R.; Mazuruk, Krysztalof;
4 APPLICANT: Tombar-Truk, Joyce
5 TITLE OF INVENTION: FISHING EQUIPMENT
6 TITLE OF INVENTION: BIRCHEN FISHING CHARACTERIZATION SYSTEM
7 TITLE OF INVENTION: ORGANIZATION AND SUPPORT OF THE FISH FARM
8 NUMBER OF SEQUENCES: 48
9 ADDRESS WHERE ADDRESS:
10 ADDRESS: Morgan & Finnegan
11 STREET: 445 Park Avenue
12 CITY: New York
13 STATE: New York
14 COUNTRY: USA
15 ZIP: 10154
16 COMPUTER READABLE FORM:
17 MEDIUM TYPE: Floppy Disk
18 COMPUTER: IBM PC compatible
19 LANGUAGE: English, M. I. S
20 SOFTWARE: WORDPERFECT 5.1
21 CURRENT APPLICATION DATA:
22 APPLICATION NUMBER: 05/29/007, 041A
23 FILING DATE: 30-DEC-1994
24 CLASSIFICATION: 4.85
25 PRIOR APPLICATION DATA:
26 APPLICATION NUMBER: 08/257,963
27 FILING DATE: 07-JUN-1994
28 PRIOR APPLICATION DATA:
29 APPLICATION NUMBER: 07/052,796
30 FILING DATE: 24-SEP-1992
31 ALTERNATIVE INFORMATION:
32 NAME: Jacoby R. AUTH
33 REGISTRATION NUMBER: 46454
34 REFERENCE TO SECT NUMBER: 2026412,0052
35 INFORMATION INFORMATION:
36 TELEPHONE: (212) 758-4800
37 TELEFAX: (212) 751-6849
38 INFORMATION PER SEQ. ID NO.: 48
39 SEQUENCE CHARACTERISTICS:
40 LENGTH: 2261 base pairs
41 TYPE: Nucleic Acid
42 STRANDNESS: Double
43 TOPOLOGY: Unknown
44 MOLECULE TYPE: Genomic DNA
45 FEATURE:
46 NAME/KEY: F1 147
47 LOCATION:
48 IDENTIFICATION METHOD:
49 OTHER INFORMATION: full length sequence
50 OTHER INFORMATION: sequence for F1F1 plus flanking sequences.
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05-08-467 041A-4

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Query Match	Score (1.6)	Length (2.4)
Best Local Similarity	81.8%	Prod. No. 15045
Matches	253	Mismatches 44
		Indels 15
		Gaps 4





DB 6247 1CTGTCTGCAAAAAAAAAAAAAAAAAAAAA 6218

## RESULT 8

US-08-370-975B-1/c

Sequence 1: Application US/08370975B

Patent No. 5622851

GENERAL INFORMATION:

APPLICANT: Maley, Frank

APPLICANT: Maley, Gladys F.

APPLICANT: Maley, Kiroo Y. R.

TITLE OF INVENTION: Human Recombinant Platelet Lectinase Gene

NUMBER OF SEQUENCES: 14

CORRESPONDENCE ADDRESS:

ADDRESSEE: Nixon, Harshave, Devans A Bayle

STREET: Clinton Square, P.O. Box 1051

CITY: Rochester

STATE: New York

COUNTRY: USA

ZIP: 14603

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1 0, Version #1.05

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/370, 975B

FILING DATE: 10-JAN-1995

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Timiao, Susan J.

REGISTRATION NUMBER: 34,103

REFERENCE/DOCKET NUMBER: 20894/80

TELECOMMUNICATION INFORMATION:

TELEPHONE: (716)263-1636

TELEFAX: (716)263-1600

INFORMATION FOR SEQ ID NO: 1

SEQUENCE CHARACTERISTICS:

LENGTH: 26764 base pairs

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: LNA (genomic)

POSITION IN GENOME:

CHROMOSOME/SEGMENT: 4q35

US-08-370-975B-1

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

Matches 959; Consistency 9; Mismatches 68; Indels 12; Cys 2;

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

Matches 959; Consistency 9; Mismatches 68; Indels 12; Cys 2;

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

Matches 959; Consistency 9; Mismatches 68; Indels 12; Cys 2;

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

Matches 959; Consistency 9; Mismatches 68; Indels 12; Cys 2;

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

Matches 959; Consistency 9; Mismatches 68; Indels 12; Cys 2;

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

Matches 959; Consistency 9; Mismatches 68; Indels 12; Cys 2;

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

Matches 959; Consistency 9; Mismatches 68; Indels 12; Cys 2;

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

Matches 959; Consistency 9; Mismatches 68; Indels 12; Cys 2;

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

Matches 959; Consistency 9; Mismatches 68; Indels 12; Cys 2;

Query Match

Best local similarity 75.8%; Score 177.2; DB 1; Length 26764;

DB 8210 1CTGTCTGCAAAAAAAAAAAAAAAAAAAAA 8181

## RESULT 9

US-09-798-096-10/c

Sequence 10: Application US/09798096

Patent No. 6499378

GENERAL INFORMATION:

APPLICANT: Donna L. Ward

APPLICANT: Andrew T. Watt

TITLE OF INVENTION: ANTISENSE REGULATION OF PROTEIN EXPRESSION

FILE NUMBER: 6499378

CURRENT FILING DATE: 2001-02-01

NUMBER OF SEQ ID NOS: 89

SEQ ID NO 10

LENGTH: 96500

TYPE: LNA

ORGANISM: Homo sapiens

FEATURES:

US-09-798-096-10

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;

Query Match

Best local similarity 75.8%; Score 174.4; DB 4; Length 96500;

Matches 247; Consistency 9; Mismatches 66; Indels 13; Cys 2;













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